

# Alignment View Print

	1	2	3	4	5	6	7	8	9	10		20	21	22	23	24	25	26	27	28	29	30
seq2 seq	M	E																				
seq26	M	E																				
Consensus	M	E																				
	40	41	42	43	44	45	46	47	48	49		50	51	52	53	54	55	56	57	58	59	60
seq2 seq	G	A																				
seq26	G	A																				
Consensus	G	A																				
	70	71	72	73	74	75	76	77	78	79		80	81	82	83	84	85	86	87	88	89	90
seq2 seq	R	V	K	E	A	G	G	E	L													
seq26	R	V	K	E	A	G	G	E	L													
Consensus	R	V	K	E	A	G	G	E	L													
	100	101	102	103	104	105	106	107	108	109		110	111	112	113	114	115	116	117	118	119	120
seq2 seq	K	C	L	C	C	C	M	C	D													
seq26	K	C	L	C	C	C	M	C	D													
Consensus	K	C	L	C	C	C	M	C	D													
	130	131	132	133	134	135	136	137	138	139		140	141	142	143	144	145	146	147	148	149	150
seq2 seq	C																					
seq26	C																					
Consensus	C																					
	160	161	162	163	164	165	166	167	168	169		170	171	172	173	174	175	176	177	178	179	180
seq2 seq	C	K	G	G																		
seq26	C	K	G	G																		
Consensus	C	K	G	G																		
	190	191	192	193	194	195	196	197	198	199		200	201	202	203	204	205	206	207	208	209	210
seq2 seq	H	C																				
seq26	H	C																				
Consensus	H	C																				

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# Alignment View Print

	220	230	240
seq2 seq	V L V L V C V C	L T C V K R R K	R C D V V K V V
seq26	V L V L V C V C	L T C V K R R K	R C D V V K V V
Consensus	V L V L V C V C	L T C V K R R K	R C D V V K V V
	250	260	270
seq2 seq	V C R K R C E A E S	E A V I E A L O A	D V V A V E
seq26	V C R K R C E A E S	E A V I E A L O A	D V V A V E
Consensus	V C R K R C E A E S	E A V I E A L O A	D V V A V E
	280	290	300
seq2 seq	E T T T T G R	N H	
seq26	E T T T T G R	N H	
Consensus	E T T T T G R	N H	

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